

Type 2 Diabetes–Associated Hepatocellular Carcinoma: A Molecular Profile

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Hepatocellular carcinoma (HCC) is one of the leading causes of cancer-related deaths worldwide. HCC develops mainly on a background of chronic liver disease, with an incidence rate of 2% to 4% per year in cirrhotic patients. The main causative agents of the underlying liver disease and HCC are viral infections by hepatitis B virus (HBV) and hepatitis C virus (HCV), as well as alcohol abuse. Recently, new risk factors for HCC have emerged in developed countries, including nonalcoholic fatty liver disease (NAFLD) caused by metabolic syndrome (MS). Among the different components of the MS, type 2 diabetes, characterized by hyperglycemia, hyperinsulinemia,

and insulin resistance, has been identified as an independent risk factor for HCC. The aim of this brief review is to describe the impact of insulin resistance and type 2 diabetes on the incidence of HCC, the underlying molecular defect linked to tumor development, and potential future therapeutic strategies identified by analysis of epidemiological studies and *in vitro* experiments.

EPIDEMIOLOGICAL EVIDENCE

Epidemiological studies have revealed that type 2 diabetes plays an independent role in HCC development (Table 1). Diabetes has been associated with a 2- to 3-

Abbreviations: AOR, adjusted odds ratio; CI, confidence interval; HBV, hepatitis viral B; HCC, hepatocellular carcinoma; HCV, hepatitis viral C; HR, hazard ratio; IGF, insulin growth factor; IGFBP, IGF binding protein; IGF1R, IGF1 receptor; IL-6, interleukin-6; JNK1, c-jun amino terminal kinase 1; LPS, lipopolysaccharide; MAPK, mitogen-activated protein kinase; MS, metabolic syndrome; mTOR, mammalian target of rapamycin; NAFLD, nonalcoholic fatty liver disease; NF-κB, nuclear factor-κB; OR, odds ratio; ROS, reactive oxygen species; TNF-α, tumor necrosis factor-α; VEGF, vascular endothelial growth factor.

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TABLE 1. RISK FOR HCC AND TYPE 2 DIABETES

Authors	Association	Description of the Study
Fu et al. ¹¹	HR 1798 (95% CI 1.194-2.707)	Cohort study
Aliment Pharmacol Therapy 2015		New-onset diabetes, patients with HBV
Raff et al. 12	HR 3.0 (95% CI 1.3-6.9)	Cohort study
J Clin Transl Hepatol 2015		Patients with NAFLD or alcoholic liver disease
Setiawan et al. 13	RR 2.62 (95% CI 2.13-3.23)	Cohort study
J Natl Cancer Inst 2014		Increased risk in all ethnic groups
Arase et al. 14	HR 1.73 (95% CI 1.3-2.3)	Cohort study
Hepatology 2013		Patients with HCV
Turati et al. ¹⁵	OR 4.33 (95% CI 1.89-9.86)	Case-control study
Br J Cancer 2013		
Koh et al. 16	HR 2.14 (95% CI 1.69-2.71)	Cohort study
Br J cancer 2013		
Schlesinger et al. 17	RR 2.17 (95% CI 1.36-3.47)	Cohort study
Ann Oncol 2013		
Atchison et al. 18	RR 1.95 (95% CI 1.82-20.9)	Cohort study
Int J Cancer 2011		
Hassan et al. 19	AOR 4.2 (95% CI 3.0-5.9)	Case-control study
Cancer 2010		Decreased risk for HCC using metformin
Veldt BJ et al. ²⁰	HR 3.28 (95% CI 1.35-7.97)	Cohort study
Hepatology 2008		Patients with HCV
Rousseau et al. ²¹	OR 3.1 (95% CI 1.1-8.8)	Case-control study
Int J cancer 2006		
Davila et al. ²²	OR 3.8 (95% CI 2.74-3.46)	Case-control study
Gut 2005		Increased risk regardless of other HCC risk factors
El-Serag et al. ²³	HR 2.16 (95% CI 1.86-2.52)	Cohort study
Gastroenterology 2004		Increased risk for chronic liver diseases
Hassan et al. ²⁴	OR 4.3 (95% CI 1.9-9.9)	Case-control study
Hepatology 2002		Synergism with alcohol and viral hepatitis

AOR, adjusted odds ratio; CI, confidence interval; HR, hazard ratio.

fold increase in the risk for HCC occurrence; this risk did not differ between ethnic populations in the United States. The risk is even higher with duration of diabetes, with an odds ratio (OR) of 2.2 in patients having a longer than 10-year duration (Table 1). New-onset diabetes has also been significantly associated with higher incidence of HCC. Strong evidence also describes a synergistic interaction between type 2 diabetes and other risk factors for HCC, such as chronic HCV and HBV infection and alcohol abuse. In a prospective cohort of patients with HCV-related cirrhosis, insulin resistance was also an independent risk factor for HCC development. Finally, a subset of HCC related to MS developed on noncirrhotic liver, suggesting a direct oncogenic mechanism. ²

PHYSIOPATHOLOGY AND MOLECULAR PATHWAYS

In cellulo models, as well as mouse models, have helped to dissect the molecular pathway involved in liver carcinogenesis caused by insulin resistance and obesity (Fig. 1).

Hyperinsulinemia and obesity promote chronic liver inflammation and steatosis by the release of proinflammatory cytokines such as tumor necrosis factor- α (TNF- α), interleukin-6 (IL-6), and nuclear factor- κ B (NF- κ B) (Fig. 1). This inflammatory state is a conjunction of the action of hepatocytes, Kupffer cells, and adipocytes. Moreover, MS increases the circulating level of leptin, a proinflammatory cytokine, but also decreases the

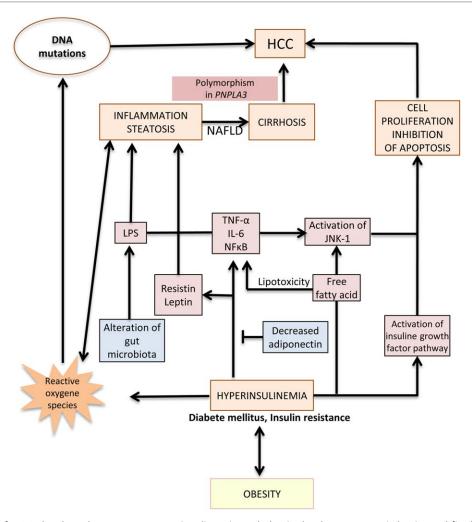


FIG 1 Pathogenesis of HCC developed on NAFLD. Hyperinsulinemia and obesity lead to NAFLD, cirrhosis, and finally, HCC development. However, a subset of NAFLD-related HCC could also develop on noncirrhotic liver. Chronic liver inflammation, alteration of gut microbiota, lipotoxicity of free fatty acid, production of ROS, and activation of the IGF pathway and the JNK1 pathway promote NAFLD and HCC development. A polymorphism of *PNPLA3* (Patatin-like phospholipase domain containing 3, rs738409 c444C>G, I148M minor allele) increased the risk for NAFLD-related cirrhosis and NAFLD-related HCC. Abbreviation: LPS, lipopolysaccharide.

production of adiponectin, an anti-inflammatory polypeptide that also inhibits angiogenesis in animal models. Hyperglycemia leads to free fatty acid release that induces accumulation of reactive oxygen species (ROS) (Fig. 1). ROS promote carcinogenesis through uncontrolled liver inflammation, steatosis, and cell proliferation, and may induce cancer-promoting mutations in tumor suppressor gene *TP53*. Moreover, free fatty acid production activates c-jun amino terminal kinase 1 (JNK1) that induces cellular proliferation and inhibition of apoptosis. ⁴

In human HCC, a subset of tumors have been shown to harbor activation of the insulin growth factor (IGF) pathway.⁶ IGF1 induces phosphorylation of insulin receptor substrate 1, leading to activation of AKT/mammalian

target of rapamycin (MTOR) and mitogen-activated protein kinase (MAPK) pathways that inhibit apoptosis and promote cell proliferation (Fig. 2). Upregulation of the IGF pathway by hyperinsulinemia is the consequence of overexpression of ligands such as IGF1 and aberrant expression of fetal IGF2. Concomitant downregulation of negative regulators of the pathway, such as the suppressor of cytokine signaling protein, is also observed in tumors, along with decreased production of IGF binding protein (IGFBP) type 1 and IGFBP2 that increases IGF1 bioavailability (Fig. 2). Consequently, the IGF axis is a key signaling pathway involved in liver carcinogenesis related to type 2 diabetes and obesity. Moreover, each human HCC is a unique combination

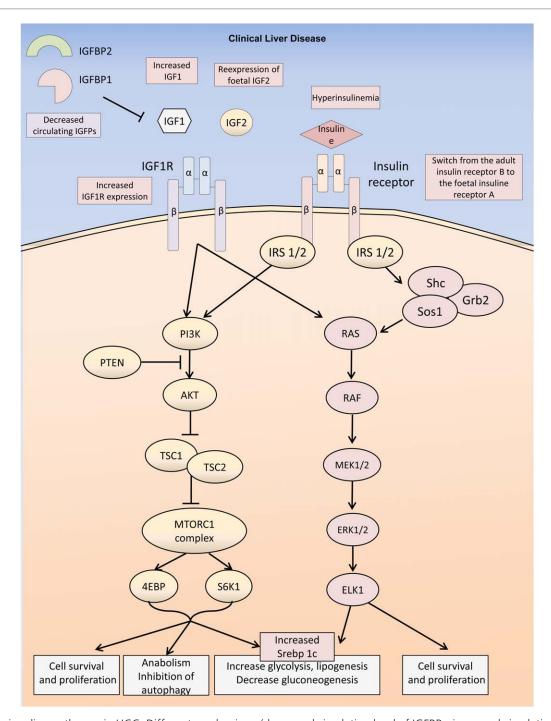


FIG 2 Insulin signaling pathways in HCC. Different mechanisms (decreased circulating level of IGFBPs, increased circulating level of IGF1, reexpression of the foetal IGF2, overexpression of IGF1R, hyperinsulinemia, reexpression of the foetal form of the insulin receptor (IR) type A instead of the adult form IRB) lead to the activation of the IGF1 receptor (IGF1R) and insulin receptor signaling pathways. Most of the metabolic and proliferative consequences are mediated through the downstream activation of Akt/Mtor and ras/raf/map kinase signaling.

of somatic genetic alterations, with a mean number of 40 to 60 mutations per tumor in the coding sequence.⁸ Somatic mutations in driver genes were also closely related to risk factors and etiology: *TP53* mutations with HBV infection, the *TP53* R249S mutation with

aflatoxin B1 exposure, and *ARID1A* and *CTNNB1* mutations with alcohol consumption.⁸ However, no clear association between genetic defects or nucleotide signatures and type 2 diabetes and MS have thus far been identified.

INSULIN RESISTANCE AND LIVER CARCINOGENESIS: A NEW THERAPEUTIC STRATEGY

Human epidemiological studies have shown that statins confer protection against development of steatohepatitis, fibrosis, and HCC, mainly in patients with diabetes and obesity. El-Serag et al.⁹ compared 1303 patients with diabetes and 5212 control subjects, and identified a 25% reduction in the risk for development of HCC in patients with diabetes using statins. Statins have no direct effect on insulin resistance, but the antitumor action could be explained by an anti-inflammatory property mediated by inhibition of the JNK and MAPK pathways.

Moreover, in epidemiological studies performed on a general population, HBV- or HCV-infected patients showed that metformin intake was also associated with a reduced incidence of HCC in a time- and dose-dependent manner. Metabolic and antiangiogenic effects have been proposed as potential antitumor mechanisms. Metformin activated AMP-activated protein kinase that inhibited the mTOR complex and fostered cell cycle arrest *in vitro*. Metformin also exhibited antiangiogenic effects through downregulation of vascular endothelial growth factor. Moreover, different signaling pathways such as Wnt/β-catenin or transcription factors such as Myc proto-oncogene protein were shut down by metformin *in cellulo*.

Finally, an anti-IGF1 receptor (anti-IGF1R) antibody, cixutumumab, has been tested in patients with advanced HCC, as well as a small tyrosine kinase inhibitor, OSI-906, targeting IGF1R/IR. However, they demonstrated no antitumor activity and have metabolic toxicities. An antibody against IGF2 is currently tested in solid cancers including HCC.⁷

In conclusion, type 2 diabetes and obesity are key players in NAFLD and HCC development. Chronic liver inflammation, modulation of the immune response, lipotoxicity, production of ROS, and activation of the IGF pathway are some of the mechanisms that promote tumorigenesis in this setting. Finally, epidemiological and *in vitro* studies suggested a chemopreventive effect of statin and metformin on HCC development. This will require confirmation by randomized controlled trials.

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